

LIST OF REFERENCES CITED BY APPLICANT <i>(Use several sheets if necessary)</i>				ATTY. DOCKET NO.		APPLICATION NO.	
				9882-015-999		09/805,353	
				APPLICANT			
				Gonzalez et al.			
				FILING DATE		GROUP	
				March 13, 2001		TBB 1631	
U.S. PATENT DOCUMENTS							
*EXAMINER INITIAL		DOCUMENT NUMBER	DATE	NAME	CLASS	SUBCLASS	FILING DATE IF APPROPRIATE
	AA						
	AB						
FOREIGN PATENT DOCUMENTS							
		DOCUMENT NUMBER	DATE	COUNTRY	CLASS	SUBCLASS	TRANSLATION
							YES NO
	AC						
	AD						
OTHER REFERENCES (Including Author, Title, Date, Pertinent Pages, Etc.)							
	AE	Barber, Ken, 1999, Signals Online Magazine.					
<i>ml</i>	AF	Baxter et al., 1998, Proteins: Structure, Function and Genetics 33:367-382.					
<i>ml</i>	AG	Böhm, Hans-Joachim, , 1998, J. of Computer-Aided Molecular Design, 12: 309-323.					
	AH	Bower et al., 1997, www.empharm, ucwf.edu					
<i>ml</i>	AI	Craven et al., 1998, Nucleic Acids Research, 26: 21:5007-5008					
<i>ml</i>	AJ	Dahiyat, B.I. & Mayo, S.L. De novo protein design: fully automated sequence selection. Science 278, 82-87. (1997)					
<i>ml</i>	AK	Debouck et al., 1999, Nature Genetics Supplement, 21:48-50.					
	AL	Dunbrack, 1999, Backbone-dependent rotamer library webpage					
<i>ml</i>	AM	Dunbrack, 1997, Protein Science, 6: 1661-1681.					
<i>ml</i>	AN	Eldridge et al., 1997, J. of Computer-Aided Molecular Design, 11: 425-445.					
<i>ml</i>	AO	Emili et al, 2000, Nature Biotechnology, 18: 393-397.					
<i>ml</i>	AP	Gee, S.H., Quenneville, S., Lombardo, C.R. & Chabot, J. Single-amino acid substitutions alter the specificity and affinity of PDZ domains for their ligands. Biochemistry 39, 14638-14646. (2000)					
	AQ	Gerhold et al., 1999, TIBS, p. 168-173					
	AR	Guex et al., DEA-SIB: Module 6, www.expasy.ch/swissmod/course,					
<i>ml</i>	AS	Jarvik et al., Annu. Rev. Genet., 1998, 32:601-18					
<i>ml</i>	AT	Kauvar et al., 1995, Chemistry & Biology, 2:107-118					
<i>ml</i>	AU	Lennon et al., 1991, TIG 7:,10: 314-317.					
<i>ml</i>	AV	Levchenko, I., Smith, C.K., Walsh, N.P., Sauer, R.T. & Baker, T.A. PDZ-like domains mediate binding specificity in the Ckp/Hsp 100 family of chaperones and protease regulatory subunits. Cell 91, 939-947. (1997)					
<i>ml</i>	AW	Lueking et al., 1999, Analytical Biochemistry 270:103-111.					
<i>ml</i>	AX	MacBeath et al., 2000, Science, 289:1760-1763.					
	AY	Martz et al., Protein Data Bank, www.rcsb.org/pdb/experimental_methods.html					
<i>ml</i>	AZ	Martzen et al., 1999, 286:11531155					

<i>ml</i>	BA	Murray et al., 1998, J. of Computer-Aided Molecular Design, 12:503-519.
<i>ml</i>	BB	Oldenburg et al., 1997, Nucleic Acids Research, 1997, 25:2:451-452.
<i>ml</i>	BC	Patel et al., 1998, J. of computer-Aided.Molecular Design, 12:543-556.
	BD	Riechmann et al, 1992, J. Molecular Biology, 913-918
<i>ml</i>	BE	Scheer et al, 1997, Proc. National Academy of Sciences, 94:808-813.
<i>ml</i>	BF	Schneider, S. et al. Mutagenesis and selection of PDZ domains that bind new protein targets. Nat Biotechnol 17, 170-175. (1999).
<i>ml</i>	BG	Skerra et al., 1999, Elsevier Biomolecular Engineering 16:79-86.
<i>ml</i>	BH	Uetz et al, 2000, Nature 403:623-631.
<i>ml</i>	BI	Walter et al., 2000, Current Opinion in Microbiology, 3: 298-302
<i>ml</i>	BJ	Warshel et al., 1986, Proc. Natl. Academy of Science, 83:3806-3810.
	BK	www.cmbi. Kun.nl/whatif/
	BL	EMBL-EBK MaxSprout server - http://www2.embl.ebi.ac.uk/dali/maxsprout/
	BM	MicroCal - http://www.microcalorimetry.com/
	BN	NCBI Structure- http://www.ncbi.nlm.nih.gov/Structure/mmdb/mdb.shtml
	BO	http://www.ncbi.nlm.nih.gov
	BP	Structural Bioinformatics - http://www.rcsb.org/index.html
	BQ	(Various authors) 1991-1999, Protein-Protein Interaction Abstracts
EXAMINER <i>[Signature]</i>		DATE CONSIDERED 7/15/2004
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